#Z 03-09-0

PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/09/766,378

DATE: 03/13/2001 TIME: 23:29:13

INPUT SET: S36502.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING
2 3	(1)	General Information:
4	(1)	
5 6 7 8 9		(i) APPLICANT: Rhode, Peter R. Acevedo, Jorge Burkhardt, Martin Jiao, Jin-an Wong, Hing C.
10		/dd/ mimilia on infinition. Governo a way governo and
11 · 12 13		(ii) TITLE OF INVENTION: SOLUBLE MHC COMPLEXES AND METHODS OF USE THEREOF
14 15		(iii) NUMBER OF SEQUENCES: 38
16		(iv) CORRESPONDENCE ADDRESS:
17		(A) ADDRESSEE: Dike, Bronstein, Roberts & Cushman, LLP
18 19	•	(B) STREET: 130 Water Street (C) CITY: Boston
20		(D) STATE: MA
21		(E) COUNTRY: usa
22 23		(F) ZIP: 02109
24		(v) COMPUTER READABLE FORM:
25		(A) MEDIUM TYPE: Diskette
26		(B) COMPUTER: IBM Compatible
27		(C) OPERATING SYSTEM: DOS
28 29		(D) SOFTWARE: FastSEQ for Windows Version 2.0
30		(vi) CURRENT APPLICATION DATA:
31		(A) APPLICATION NUMBER: 09/766,378
32		(B) FILING DATE:
33		(C) CLASSIFICATION:
34		/ · · · · · · · · · · · · · · · · · · ·
35		(vii) PRIOR APPLICATION DATA:
36 37		(A) APPLICATION NUMBER: 08/960,190 (B) FILING DATE:
38		(b) Filling Dail:
39		(viii) ATTORNEY/AGENT INFORMATION:
40		(A) NAME: Corless, Peter F
41		(B) REGISTRATION NUMBER: 33,860
42		(C) REFERENCE/DOCKET NUMBER: 48002
43		
44		(ix) TELECOMMUNICATION INFORMATION:
45		(A) TELEPHONE: 617-523-3400
46		(B) TELEFAX: 617-523-6440

RAW SEQUENCE LISTING PATENT APPLICATION US/09/766,378

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47	(C) TELEX:	
48		
49		
50	(2) INFORMATION FOR SEO ID NO:1:	
51		
52	(i) SEQUENCE CHARACTERISTICS:	
53	(A) LENGTH: 8 base pairs	
54	(B) TYPE: nucleic acid	•
55	(C) STRANDEDNESS: single	
56	(D) TOPOLOGY: linear	
57	(b) Toronogi. Timear	
58	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
59	(XI) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
	CONCONTIC.	
60	CCACCATG	8
61	(0) THEODINATION FOR ONE TO NO 0	
62	(2) INFORMATION FOR SEQ ID NO:2:	
63	(1) 570	
64	(i) SEQUENCE CHARACTERISTICS:	
65	(A) LENGTH: 43 base pairs	
66	(B) TYPE: nucleic acid	
67	(C) STRANDEDNESS: single	•
68	(D) TOPOLOGY: linear	
69		
70		
71	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
72	· ·	
73	CCCCCAAGC TTCCGGGCCA CCATGGCTCT GCAGATCCCC AGC	43
74		
75	(2) INFORMATION FOR SEQ ID NO:3:	
76		
77	(i) SEQUENCE CHARACTERISTICS:	
78	(A) LENGTH: 34 base pairs	
79	(B) TYPE: nucleic acid	
80	(C) STRANDEDNESS: single	
81	(D) TOPOLOGY: linear	
82		
83	·	
84	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
85		
86	CCCCCCACTT AAGGTCCTTG GGCTGCTCAG CACC	34
87		
88	(2) INFORMATION FOR SEQ ID NO:4:	
89	•	
90	(i) SEQUENCE CHARACTERISTICS:	
91	(A) LENGTH: 37 base pairs	
92	(B) TYPE: nucleic acid	
93	(C) STRANDEDNESS: single	
94	(D) TOPOLOGY: linear	
95	(b) Totologi. Ilmout	
96	•	
97	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
98	(AI) DESCRIPTION: DESCRIPTION:	
99	CCCCCCCCA TCCCCCCAAA CTCCCAAAACC CATTOTOCC	2.5

RAW SEQUENCE LISTING PATENT APPLICATION US/09/766,378

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100		
101	/2)	
102	(2) INFORMATION FOR SEQ ID NO:5:	
103	(1)	
104	(i) SEQUENCE CHARACTERISTICS:	
105	(A) LENGTH: 32 base pairs	
106	(B) TYPE: nucleic acid	
107	(C) STRANDEDNESS: single	
108	(D) TOPOLOGY: linear	
109		·
110	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
111		
112	GCGGCGACTA GTCCACTCCA CAGTGATGGG GC	32
113		
114		
115	(2) INFORMATION FOR SEQ ID NO:6:	
116	(i) SEQUENCE CHARACTERISTICS:	
117	(A) LENGTH: 36 base pairs	
118	. (B) TYPE: nucleic acid	
119	(C) STRANDEDNESS: single	
120	(D) TOPOLOGY: linear	
121		
122	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
123		
124	GGGGGGCCA TGGCCGAAGA CGACATTGAG GCCGAC	36
125		
126		
127	(2) INFORMATION FOR SEQ ID NO:7:	
128	·	
129	(i) SEQUENCE CHARACTERISTICS:	
130	(A) LENGTH: 32 base pairs	
131	(B) TYPE: nucleic acid	
132	(C) STRANDEDNESS: single	
133	(D) TOPOLOGY: linear	
134		
135	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	•
136		
137	GCGCGACTAG TCCAGTGTTT CAGAACCGGC TC	32
138		
139		
140	(2) INFORMATION FOR SEQ ID NO:8:	
141		
142	(i) SEQUENCE CHARACTERISTICS:	
143	(A) LENGTH: 31 base pairs	
144	(B) TYPE: nucleic acid	•
145	(C) STRANDEDNESS: single	
146	(D) TOPOLOGY: linear	
147	\-,	
148	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
149	(iii) degomine becommends, bug ib no.0.	
150	GGGGGGGATA TCTCTCAGGC TGTTCACGCT G	31
151	TOTAL	31
152		•

RAW SEQUENCE LISTING PATENT APPLICATION US/09/766,378

DATE: 03/13/2001 TIME: 23:29:14

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153			INPUT SET: \$36502.raw
155 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 46 base pairs 157 (B) TYPE: nucleic acid 158 (C) STRANDENNESS: single 160 161 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: 162 163 (G) STRANDENNESS: SINGLE 164 (A) SEQUENCE DESCRIPTION: SEQ ID NO:9: 163 (A) SEQUENCE DESCRIPTION: SEQ ID NO:10: 165 (A) SEQUENCE CHARACTERISTICS: (A) LENGTH: 37 base pairs (C) STRANDENNESS: single (D) TOPOLOGY: linear (D) SEQUENCE CHARACTERISTICS: (D) TOPOLOGY: linear (D)	_	(2) INFORMATION FOR SEQ ID NO:9:	
156		(i) SECTIONCE CHARACTERISTICS.	
157 (B) TYPE: nucleic acid 158 (C) STRANDEDNESS: single 159 (D) TOPOLOGY: linear 160 161 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: 162 163 GGGGGGTTCG AAAAGTGTAC TTACGGGGGG CTGGAATCTC AGGTTC 46 164 165 166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (i) SEQUENCE CHARACTERISTICS: 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 (Xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 (Xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 207 (A) LENGTH: 42 base pairs 199 208 (I) SEQUENCE CHARACTERISTICS: 209 (I) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		The state of the s	
158			
159			•
160 161 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: 162 163 GGGGGGTTCG AAAAGTGTAC TTACGGGGGG CTGGAATCTC AGGTTC 46 164 165 166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDENNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDENNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 2101 (A) LENGTH: 42 base pairs 221 (B) TYPE: nucleic acid 222 (B) TYPE: nucleic acid 233 (C) STRANDENNESS: single 244 (D) TOPOLOGY: linear			
161 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: 162 163 164 165 166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 181 182 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE DESCRIPTION: SEQ ID NO:11: 188 189 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(2) 10102001: 111001	•
162 163 GGGGGGTTCG AAAAGTGTAC TTACGGGGGG CTGGAATCTC AGGTTC 164 165 166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(xi) SEQUENCE DESCRIPTION: SEO ID NO:9:	
163 GGGGGGTTCG AAAAGTGTAC TTACGGGGGG CTGGAATCTC AGGTTC 164 165 166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDENDESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (a) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 2101 (A) LENGTH: 42 base pairs 221 (2) INFORMATION FOR SEQ ID NO:12: 222 (B) TYPE: nucleic acid 233 (C) STRANDEDNESS: single 244 (D) TOPOLOGY: linear		, ,	
165 166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		GGGGGGTTCG AAAAGTGTAC TTACGGGGGG CTGGAATCTC AGGTTC	46
166 167 168 169 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDENNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDENNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 36 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDENNESS: single 204 (D) TOPOLOGY: linear	164		
166 167 168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (i) SEQUENCE CHARACTERISTICS: 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 211 222 (B) TYPE: nucleic acid 223 (C) STRANDEDNESS: single 234 (D) TOPOLOGY: linear	165		
168 169 170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 42 base pairs 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	166		
169 170 171 172	16:7		
170 171 172 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (i) SEQUENCE CHARACTERISTICS: (a) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs 196 197 198 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 203 (A) LENGTH: 42 base pairs 204 (B) TYPE: nucleic acid 205 (C) STRANDEDNESS: single 206 (D) TOPOLOGY: linear	168		
170 171 172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single	169		
172 (2) INFORMATION FOR SEQ ID NO:10: 173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	170		,
173 174 (i) SEQUENCE CHARACTERISTICS: 175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	171		
174 (i) SEQUENCE CHARACTERISTICS: 175 (a) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	172	(2) INFORMATION FOR SEQ ID NO:10:	
175 (A) LENGTH: 37 base pairs 176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	173		
176 (B) TYPE: nucleic acid 177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	174	(i) SEQUENCE CHARACTERISTICS:	
177 (C) STRANDEDNESS: single 178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	175	(A) LENGTH: 37 base pairs	
178 (D) TOPOLOGY: linear 179 180 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10: 181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear	176		
179 180	177		
180	178	(D) TOPOLOGY: linear	
181 182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
182 GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC 37 183 184		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
183 184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
184 185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		GGGGGGCTCG AGTATCAAAG AAGAACATGT GATCATC	37
185 (2) INFORMATION FOR SEQ ID NO:11: 186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
186 187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
187 (i) SEQUENCE CHARACTERISTICS: 188 (A) LENGTH: 36 base pairs 189 (B) TYPE: nucleic acid 190 (C) STRANDEDNESS: single 191 (D) TOPOLOGY: linear 192 193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(2) INFORMATION FOR SEQ ID NO:11:	
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193 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11: 194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(D) TOPOLOGY: Ilnear	
194 195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 36 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(vi) CENTENCE DECEPTORION GEO TO NO CO	
195 GCGGCGGGAT CCGTTCTCTG TAGTCTCTGG GAGAGG 196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(XI) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
196 197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		CCCCCCCAT CCCTTCTCTC TACTCTCTC CACACC	36
197 198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		GCGGCGGGAI CCGIICICIG IAGICICIGG GAGAGG	36
198 (2) INFORMATION FOR SEQ ID NO:12: 199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
199 200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(2) INFORMATION FOR SEC ID MO-12.	
200 (i) SEQUENCE CHARACTERISTICS: 201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(2) Intolumiton for BEQ ID NO.12:	
201 (A) LENGTH: 42 base pairs 202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear		(i) SEQUENCE CHARACTERISTICS:	
202 (B) TYPE: nucleic acid 203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
203 (C) STRANDEDNESS: single 204 (D) TOPOLOGY: linear			
204 (D) TOPOLOGY: linear			
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RAW SEQUENCE LISTING PATENT APPLICATION US/09/766,378

DATE: 03/13/2001 TIME: 23:29:14

	INPUT SET: S36502	.raw
206	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
207		
208	GATAAGAGGA AGAAGAGTAC ATGCCGATGG AACCCGGGTG AG	42
209		
210 211	(2) INFORMATION FOR GEO ID NO 12.	
211	(2) INFORMATION FOR SEQ ID NO:13:	
213	(i) SEQUENCE CHARACTERISTICS:	
214	(A) LENGTH: 43 base pairs	
215	(B) TYPE: nucleic acid	
216	(C) STRANDEDNESS: single	
217	(D) TOPOLOGY: linear	
218		
219	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
220		
221	AATTCTTCAC CCGGGTTCCA TCGGCATGTA CTCTTCTTCC TCG	43
222 223		
223 224		
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226		
227		
228		
229	(2) INFORMATION FOR SEQ ID NO:14:	
230		
231	(i) SEQUENCE CHARACTERISTICS:	
232	(A) LENGTH: 75 base pairs	
233	(B) TYPE: nucleic acid	
234 235	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
236	(b) TOPOLOGI: Tiflear	
237	(xi) SEQUENCE DESCRIPTION: SEO ID NO:14:	
238	(III) DEGOLIOE DEBONITION: DEG ID NO.11.	
239	CCCCCGCTA GCGGAGGGGG CGGAAGCGGC GGAGGGGGGG ACACCCGACC ACGTTTCCTG	60
240	TGGCAGCCTA AGAGG	75
241	·	
242		
243	(2) INFORMATION FOR SEQ ID NO:15:	
244	(+) GEOVERNOR GUARAGERA	
245 246	(i) SEQUENCE CHARACTERISTICS:	
240	(A) LENGTH: 48 base pairs (B) TYPE: nucleic acid	
248	(C) STRANDEDNESS: single	
249	(D) TOPOLOGY: linear	
250	(-,	•
251	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
252		
253	CCCCCGAAT TCCCCACTAG TCCATTCCAC TGTGAGAGGG CTTGTCAC	48
254		
255	(0) TWEEDWARTON TOD GEO	•
256 257	(2) INFORMATION FOR SEQ ID NO:16:	
257 258	(i) SEQUENCE CHARACTERISTICS:	
250	(I) DEQUENCE CHARACIERIBIICS:	

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/766,378

DATE: 03/13/2001 TIME: 23:29:15

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Original Text